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>muCRAM-2
CAGAATTCGGCACGAGGTCCTGGGGGGGGGGCGACCTACGGGTTCTCCCTCAAGAGCTAATCTCTGCC
GCCACTCGCTTAGGACCTCGGGACACCGGTCCTCGGTCCACGCCCTCCCTCAACCTCTTCCACCCCTT
CAAAGAAGGACTGTCCAGACACCACTCTAGGGCCAGAGACCTGCCCTCCACGACAGTCCGCTGGAGACA
CCCCAGACCGGAGACTGACATCGGACAGGACCCGCTCTGCTTCCACTCTCAGSGGACCTCTCTCTG
CTCCGCGCGGGGAGTGTCTGGGAGACCCAGCCGCTCTGCGCTCTCAGGGGACCTCAGCTAGG
CAGCCAGCTGGGCGCGGTAGATGGCGAGGACCCCAAGGCTCTGATGCTCTGCTGCTACACTACT
TGATCGTCCCTGGACTATCATAAGGCAATGGTCTCTGCTCAATCAAGACCACTCAAGAGTCA
GTAATAGAGTTCGAAGGCTATTTGGCTGTAAACCCCAAGAGACTACCTCTCCAGACTGGAGTG
GAAGAAGGAGGACAGGGGCTCTCTGGTCTACTACCAACAGGCTCTCCAGGTGACTTTAAAGACCCGTG
CTGAGATGATAGATTCAATATACGAATCAAAATGTTACAAAGAGTGAATGCTAGAGAGTATCGCTGIGAA
GTCCAGGCTCCGACTGAGCAAGGCCAGAACCTGCGAGAGATAAAGTCAATGCTAGAGTACGATGCCAGGATA
TGCTGTTCTGCTGTAAGTCCCACTTCTGTTATGACTGGAGGTGGTGGAGTACGATGCCAGGATA
AGAAGGAACCCAGCTCCGAGTACATCTGGTTTAAAGATGGCAAGTTTGTAGGGAATCCAAAGGC
GGCACACACACAGCTCGTACACAAATGAACACGAATCTGGAATCTGCAATTCACATGATTTCCAA
GATGACAGTGGAGATATTACTGCAAGCCCGAATCTGTCCGACACCGAGGTGCTCGGAAAGCGAA
TGCAAGTAGATGTTCTCAACATAGCGGCTCATAGCAACGCTTGTGGTGGTGGCTCTCGGATTTCTGTA
TGCGGCTTGGCCTGCTATGCTCAGAGGAAAGGCTACTTTTCAAAAGAACTTCTTCCAGAGGCGAG
TCCTGCATCTAAAGTCACTACGATGGGCGAAATGATTTTCAAGGACACAAATCTCTTATAATTTAAAGA
ATTCCAGTTTGGGCTGCCCAAAACCAAGTTGTCAATGTTAATAAATTTGTAATCTCTGCTCTTACA
CTTGCAGAGTGTAGAAATATGAAGGGGAGTTTATGATCTCTAATCTCACTCAGAGAAAT
ATTTAAGCAAAACGTTCTTCCATCACTAATTAACACCTGGCATCTTGTGTTGACCTAAAGGAAATGTC
TGCTAATATCTGGTTTGAAGGCAATGAATGTGAGTTGAGTGTGACTATATCACACTGCTGTAAGG
CTAATCCAGAGCAAGATATATAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAATTC

FIG. 1B

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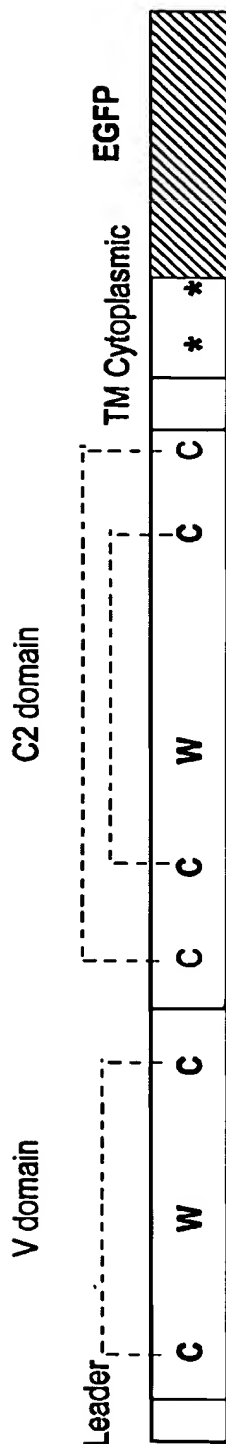
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 Docket No. 11422/264679

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JAM-EGFP
 CRAM-1-EGFP
 CRAM-2-EGFP

Tac Leader

Tac



Tac-JAM
 Tac-CRAM-1
 Tac-CRAM-2

FIG. 2A



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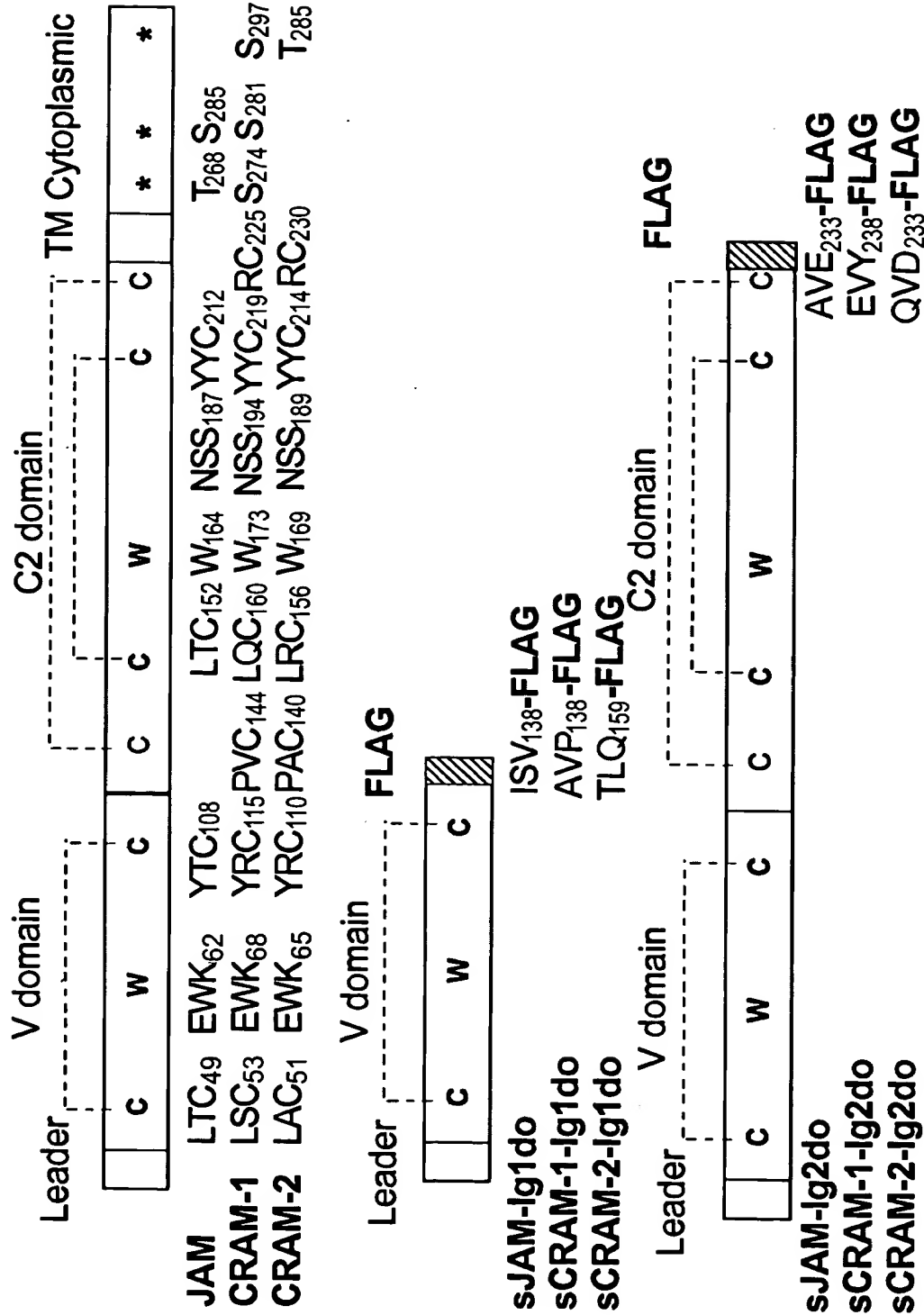


FIG. 2B

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MALSRRRLRLR LYARLPHFFL LLLFRGCMIE AVNLKSSNRN PVVHEFESVE
MARSPQGL LMLLLHYLI VALDYHKANG FSASKDHRQE VTVIEFQEAI

LSCIITHSQT SDPRIEWKKI QDGQTTYVYF DNKIQGDLAG RTDVFGKTSL
LACK-TPKKT TSSRLEWKKV GQG-VSLVYY QQALQGDFKD RAEMID-FNI

RIWNVTRSDS AIYRCEVVAL NDR-KEVDEI TIELIVQVKP VTPVCRIPAA
RIKNVTRSDA GEYRCEVSAP TEQGQNLQED KVMLEVLVAP AVPACEVPTS

VPVGKTATLQ CQSEGYPRP HYSWYRNDVP LPT-DSRANP RFQNSSFHVN
VMTGSVVELR CQDKEGNPAP EYIWFKDGTSL LG-NPKGGT H-NNSSYTNE

SETGTLVFNA VHKDDSGQYY CIASNDAGAA RCEGQDMEVY DLNIAGIIGG
HESGILQFNM ISKMDSGEYY CEARNVSGHR RCPGKRMQVD VLNISGIIAT

VLVVLIVLAV ITMGICCAVR RGCFTSSKQD GESYKSPGKH DGVNYIRTSE
VVVAVFVISV CGLGTCYAQR KGYF--SKET SFQKGSPASK -----VTTMG

EGDFRHKSSF VI
ENDFRHTKSF II

FIG. 3

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Murine and human JAM-2 (CRAM-1)

FIG. 4A

moJAM-2 (CR1)
 huJAM-2 (CR1)

10	20
<div>MA L S</div> <div>MA L R</div> <div>M A L R</div>	<div> <div>R L R L</div> <div>R L R L</div> <div>R L R L</div> </div> <div> <div>Y C</div> <div>Y C</div> <div>Y C</div> </div> <div> <div>AR L P</div> <div>AR L P</div> <div>AR L P</div> </div> <div> <div>H D</div> <div>H D</div> <div>H D</div> </div> <div> <div>FF L L L L F R</div> <div>FF L L L L F R</div> <div>FF L L L L F R</div> </div>

moJAM-2 (CR1)
 huJAM-2 (CR1)

30	40	50
<div>GC M I</div> <div>GC L I</div> <div>G C . I</div>	<div> <div>AV N L K S S N R</div> <div>AV N L K S S N R</div> <div>AV N L K S S N R</div> </div> <div> <div>T P V V</div> <div>T P V V</div> <div>T P V V</div> </div> <div> <div>H Q</div> <div>H Q</div> <div>H Q</div> </div> <div> <div>EE F E S V E</div> <div>EE F E S V E</div> <div>EE F E S V E</div> </div>	<div> <div>EE F E S V E</div> <div>EE F E S V E</div> <div>EE F E S V E</div> </div>

moJAM-2 (CR1)
 huJAM-2 (CR1)

60	70
<div>LS C I I T</div> <div>LS C I I T</div> <div>L S C I I T</div>	<div> <div>S Q T S D P P R I E</div> <div>S Q T S D P P R I E</div> <div>S Q T S D P P R I E</div> </div> <div> <div>WK K I Q D</div> <div>WK K I Q D</div> <div>WK K I Q D</div> </div> <div> <div>G E</div> <div>G E</div> <div>G E</div> </div> <div> <div>QT</div> <div>QT</div> <div>QT</div> </div>

moJAM-2 (CR1)
 huJAM-2 (CR1)

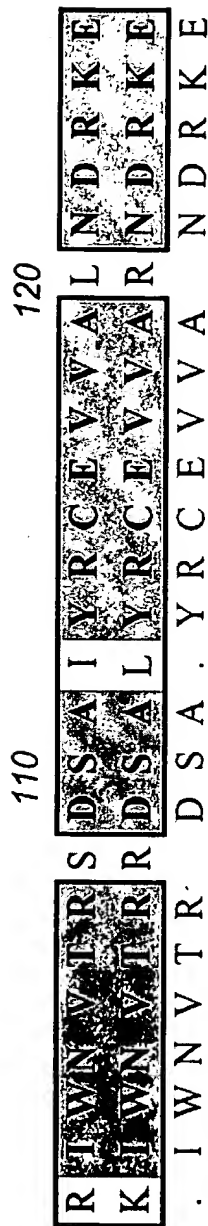
80	90	100
<div>TY V Y</div> <div>TY V F</div> <div>T Y V .</div>	<div> <div>ED N K I Q G D L A G R</div> <div>ED N K I Q G D L A G R</div> <div>ED N K I Q G D L A G R</div> </div> <div> <div>T D V</div> <div>T D V</div> <div>T D V</div> </div> <div> <div>F L</div> <div>F L</div> <div>F L</div> </div> <div> <div>G K T S L</div> <div>G K T S L</div> <div>G K T S L</div> </div>	<div> <div>G K T S L</div> <div>G K T S L</div> <div>G K T S L</div> </div>



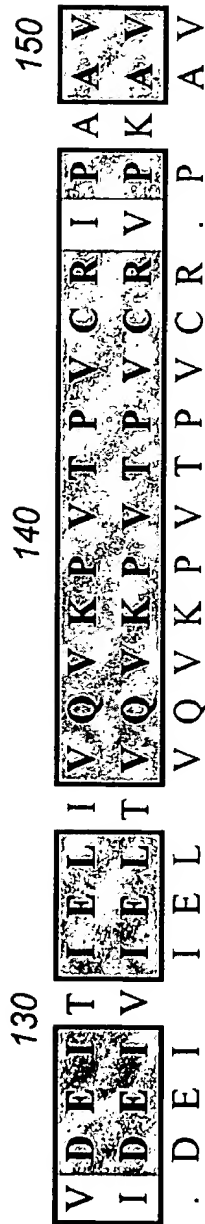
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FIG. 4B Murine and human JAM-2 (CRAM-1)

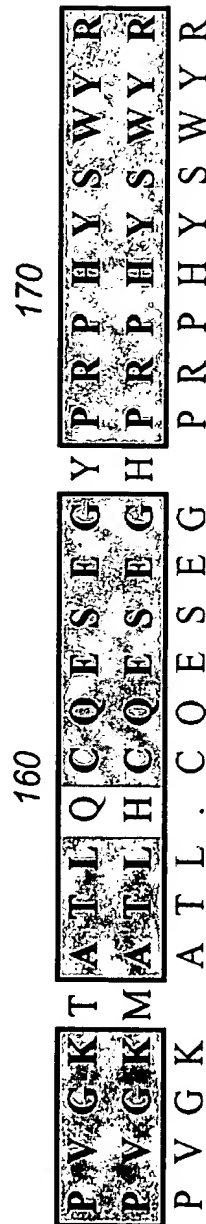
moJAM-2 (CR1)
huJAM-2 (CR1)



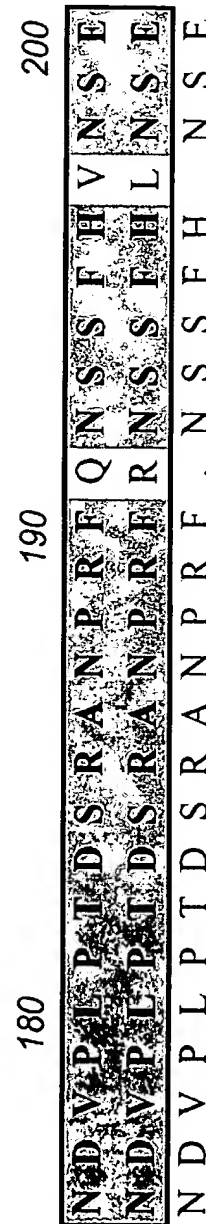
moJAM-2 (CR1)
huJAM-2 (CR1)



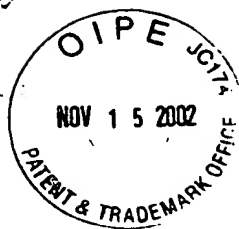
moJAM-2 (CR1)
huJAM-2 (CR1)



moJAM-2 (CR1)
huJAM-2 (CR1)



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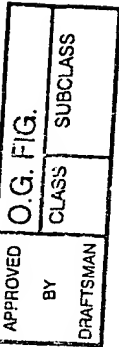
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FIG. 4C Murine and human JAM-2 (CRAM-1)

moJAM-2 (CR1)	huJAM-2 (CR1)	210		220	
		TGT L V F . A V H K D D S G Q Y Y C I A S N D A		TGT L V F . A V H K D D S G Q Y Y C I A S N D A	
moJAM-2 (CR1)	huJAM-2 (CR1)	230		240	
		G A A R C E G Q D M E V Y D L N I A G I I G G V L		G S A R C E Q E M E V Y D L N I G G I I G G V L	
moJAM-2 (CR1)	huJAM-2 (CR1)	260		270	
		V V L I V L A V I L T M G I C C A Y R R G C F I S S		V V L A V L A L I L T L G I C C A Y R R G Y F I N N	
moJAM-2 (CR1)	huJAM-2 (CR1)	280		290	
		K Q D G E S Y K S P G K H D G V N Y I R T S E E G		K Q D G E S Y K N P G K P D G V N Y I R T D E E G	
moJAM-2 (CR1)	huJAM-2 (CR1)	300		310	
		K Q D G E S Y K P G K D G V N Y I R T E E G		K Q D G E S Y K P G K D G V N Y I R T E E G	



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FIG. 4D

moJAM-2 (CR1)
huJAM-2 (CR1)

310

DTREKSSFTVI
DTREKSSFTVI

DFRHKSSFVI

330

340

350

moJAM-2 (CR1)
huJAM-2 (CR1)

360

370

moJAM-2 (CR1)
huJAM-2 (CR1)

380

390

400

moJAM-2 (CR1)
huJAM-2 (CR1)

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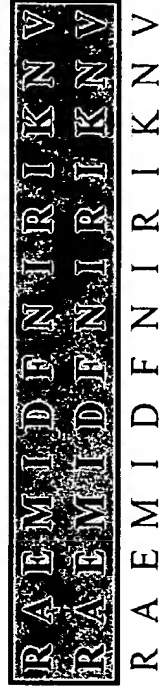
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FIG. 5A Murine and human CRAM-2 (JAM-3)

huJAM-3(CRAM-2)																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																				</
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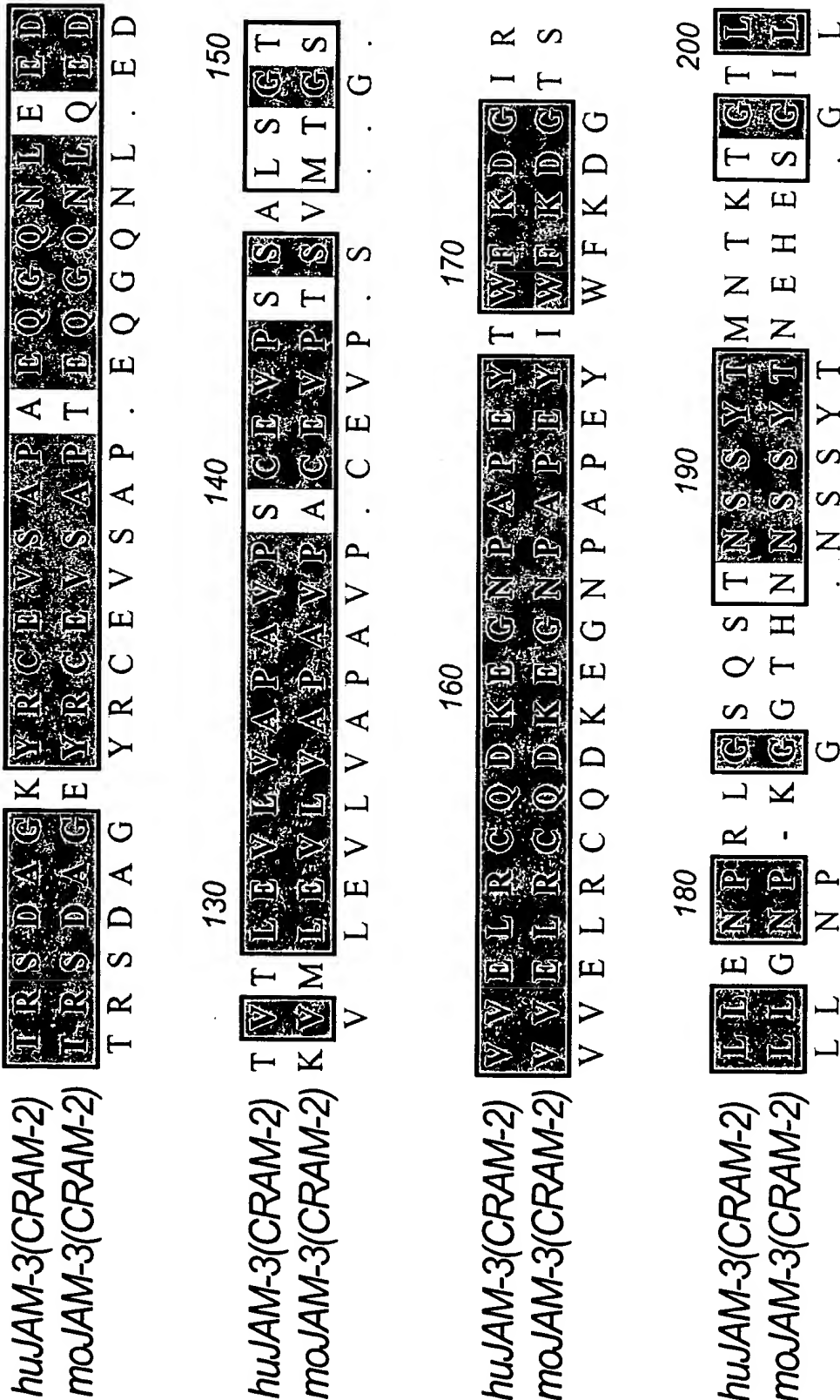
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Murine and human CRAM-2 (JAM-3)

FIG. 5B



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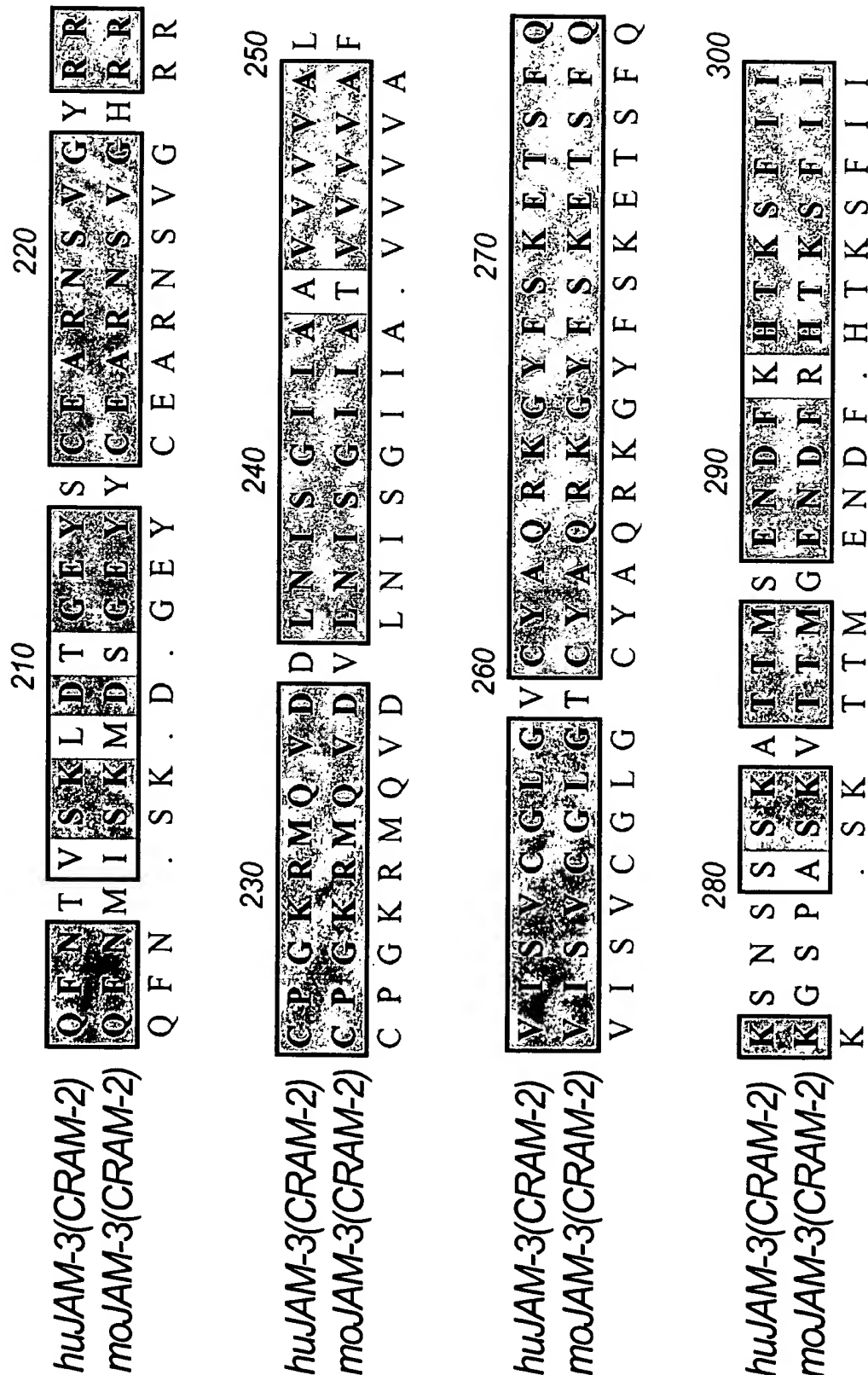
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FIG. 5C

Murine and human CRAM-2 (JAM-3)





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	CLASS	

>Nucleic acid sequence huCRAM-1/ huJAM-2

ATGGCGCTGAGGCGGCCACCGCGACTCCGGCTCTGCGCTCGGC
TGCCCTGACTTCTTCCTGCTGCTGCTTTTCAGGGGCTGCCTGAT
AGGGGCTGTAAATCTCAAATCCAGCAATCGAACCCCAAGTGGTA
CAGGAATTTGAAAGTGTGGAAGTGTCTTGCATCATTACGGATT
CGCAGACAAGTGACCCCAAGGATCGAGTGGAAGAAAATTCAAGA
TGAACAAACACATATGTGTTTTTTTGACAACAAAATTCAGGGA
GACTTGGCGGGTCTGTCAGAAATACTGGGGAAGACATCCCTGA
AGATCTGGAATGTGACACGGAGAGACTCAGCCCTTTATCGCTG
TGAGGTCGTTGCTCGAAATGACCGCAAGGAAATTGATGAGATT
GTGATCGAGTTAACTGTGCAAGTGAAGCCAGTGACCCCTGTCT
GTAGAGTGCCGAAGGCTGTACCAAGTAGGCAAGATGGCAACACT
GCACTGCCAGGAGAGTGAGGGCCACCCCGGCCTCACTACAGC
TGGTATCGCAATGATGTACCACTGCCACAGGATTCCAGAGCCA
ATCCAGATTTTCGCAATTCTTCTTTCCACTTAACTCTGAAAC
AGGCACTTTGGTGTTCACTGCTGTTTCAAGGACGACTCTGGG
CAGTACTACTGCATTGCTTCCAATGACGCAGGCTCAGCCAGGT
GTGAGGAGCAGGAGATGGAAGTCTATGACCTGAACATTGGCGG
AATTATTGGGGGGGTTCTGTTGTCTTGTCTGTACTGGCCCTG
ATCACGTTGGGCATCTGCTGTGCATACAGACGTGGCTACTTCA
TCAACAATAAACAGGATGGAGAAAGTTACAAGAACCCAGGGAA
ACCAGATGGAGTTAACTACATCCGCACTGACGAGGAGGGCGAC
TTCAGACACAAGTCATCGTTTGTGATCTGAGACCCGGGTGTGG
CTGAGAGCGCACAGAGCCGCACGTGCACATACCTCTGCTAGAA
ACTCCTGTCAAGGCAGCGAGAGCTGATGCACTCGACAGAGCTA
GACACTCTTCAAAGCTTTTCGTTTGGCAAGGTGACCACTACTC
TTTTACTCTACAAGCCCATGAAAAGAGAAATTTTCTCAAGAGG
ACCCGGAATATAACCCCAAGGAACCAAATGGGTGCGTTTAC
TGAGGTGGGGTCCTTAATTTGTTTTTGGCCTGATTCCTCATGAA
AATAAGGGGTCTTTAAGAGTTTGGTACGTAAAACCCCCCGCTT
GGGCCTTGGAACACATGTTTACCACCTGCGTTAAAAAAA
AAAAAA

FIG. 6A



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>huCRAM-1/huJAM-2 complete

MALRRPPRLRLCARLPDFFLLLLFRGCLIGAVNLKSSNRTPVV
QEFESVELSCIITDSQSDPRIEWKKIQDEQTTYVFFDNKIQG
DLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEI
VIELTVQVKPVTVPVCRVPKAVPVGKMATLHCQESEGHPRPHYS
WYRNDVPLPTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSG
QYYCIASNDAGSARCEEQEMEVDLNIIGGIIGGVLVVLAVLAL
ITLGICCAYYRRGYFINNKQDGESYKNPGKPDGVNYIRTDEEGD
FRHKSSFVI

>huCRAM-2/huJAM-3 partial

RAEMIDFNIRIKNVTRSDAGKYRCEVSAPAEQOGONLEEDTVTL
EVLVAPAVPSCEVPSSALSGTVVELRCQDKEGNPAPEYTWFKD
GIRLLENPRLGSOSTNSSYTMNTKTGTLOFNTVSKLDTGEYSC
EARNVSGYRRCPGKRMQVDDLNIISGIIAAVVVVALVISVCGLG
VCYAQRKGYFSKETSFQKSNSSSKATTMSENDFKHTKSFI

FIG. 6B



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APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 7B

t-end
B16
t-end/B16



CRAM-1



FIG. 7A

Y R C X A S1
TAY AGN TGY NNN GCY TCY AA

Y R C X A S2
TAY AGN TGY NNN GCY AGY AA

Y Q C X A S1
TAY CRG TGY NNN GCY TCY AA

Y Q C X A S2
TAY CRG TGY NNN GCY AGY AA

Y Y C X A S1
TAY TAY TGY NNN GCY TCY AA

Y Y C X A S2
TAY TAY TGY NNN GCY AGY AA

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		



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gacattcccctcgacatggcgctgagccggcgctgcgacttcgactgtacgcgcgctg 60
 M A L S R R L R L R L Y A R L 15
 cctcacttcttctgctgcttctcagggctgcatgatagaggcagtgaaatctcaaa 120
 P H F L L L L F R G C M I E A V N L K 35
 tccagcaaccgaaaccagtggtacatgaatttgaaagtgtggaattgtcttgcatt 180
 S S N R N P V V H E F E S V E L S (C) I I 55
 acgcactcacagacaagtgaccctaggattgaatggaagaaatccaagatggccaaacc 240
 T H S Q T S D P R I E W K I Q D G Q T 75
 acatatgtgtatttgacacaagaattcaaggagacctggcaggtcgacagatgtgttt 300
 T Y V Y F D N K I Q G D L A G R T D V F 95
 ggaaaaacttccctgaggatctggaatgtgacacgatcggatttcagccatctatcgctgt 360
 G K T S L R I W N V T R S D S A I Y R (C) 115
 gaggtcgctctaataatgaccgaaagaagtgtgatgagattaccattgagttaattgtg 420
 E V V A L N D R K E V D E I T I E L I V 135
 caagtgaagccagtgaccctgtctgcagaattccagccgctgtacctgtaggcaagacg 480
 Q V K P V T P V C R I P A A V P V G K T 155

FIG. 8A



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APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

gcaacactgcagtccaagagagcagggtatccccggcctcactacagctggtaaccgc 540
A T L Q (C) Q E S E G Y P R P H Y S W Y R 175
aatgatgtgccactgcctacagattccagagccaatcccagggtccagaattcctcttc 600
N D V P L P T D S R A N P R F Q N S S F 195
catgtgaactcggagacaggcactctggttttcaatgctgtccacaaggacgactctggg 660
H V N S E T G T L V F N A V H K D D S G 215
cagtactactgcattgcttccaatgacgcagggtgcagccagggtgtgaggggcaggacatg 720
Q Y Y (C) I A S N D A G A R C E G Q D M 235
gaagtctatgattgaacattgctgggattattggggagtccttgtgtccttattgtt 780
E V Y D L N I A G I I G G V L V L I V 255
cttgctgtgattacgatgggcatctgtgtgcgtacagacgaggctgttcatcagcagt 840
L A V I T M G I C C A Y R R G C F I S S 275
aaacaagatggagaaagctataagagcccagggaagcatgacggtgttaactacatccgg 900
K Q D G E S Y K S P G K H D G V N Y I R 295
acgagtgaggagggtgacttcagacacaaaatcgtccttgttatctgacacctgtcggct 960
T S E E G D F R H K S S F V I * 310

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FIG. 8B



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APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

gggagagacatgcaagtacctctgttggaagctggtcacagggtgctgtgagccccaga 1020
gctcctgacaaagcccccgggcagaagcttttgttttgccaaagtgtgatgactcctt 1080
ccttccttccttcctttaacaagccacaagaataaaagggaagcctcctgaagatggat 1140
gtagacacagattgttgctagcctgacctcattatggggtatagggtgatcttcaaggcc 1200
tttctggtcctcctcccatgagggaatttggaactgttttgccccaggctgttta 1260
gctgccaggacaactggcagagagaggtgagggcgtgggctgcagtagcagcaggca 1320
acagcctgatgcctgtgacagtgtcccccaggagggttttcaggcagtgcccttgctccctgg 1380
accctgacccaccgtgtgcctctgttgattggccagtagtgcatttccatcctggaga 1440
atgtgtttggaatcagcattttataaaaaaccacaatcagaaagggtgaaattgcttgctg 1500
ggaagagggtctgacccaggaaactctcctcccaagagatgccaggagataggagaac 1560
ctgtctgtcttaagtctgaaatgtgactgaagtctcctttctattgggtcttgcttattt 1620
tataaaaaatttaacattctaaattttgctagagatgtattttgattactgaaaaatttcta 1680
tataaactgtaaatatattgccatacagtggttcaaaaaagttatttttataatgagttc 1740
aactaagggtagaaggcttgggctgctagtgtttaattggaaaaataccagtagtaagtc 1800
ttttaaggagttttcttaaggagggtggctgaaatatcctttgttcaaaagaagttag 1860
catttttcataagaaaaacttactctgtctgaccactgttgcttaggaaaccattaaagaa 1920
ttccaatctaaaaaaaaaaaa

FIG. 8C



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APPROVED BY CRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

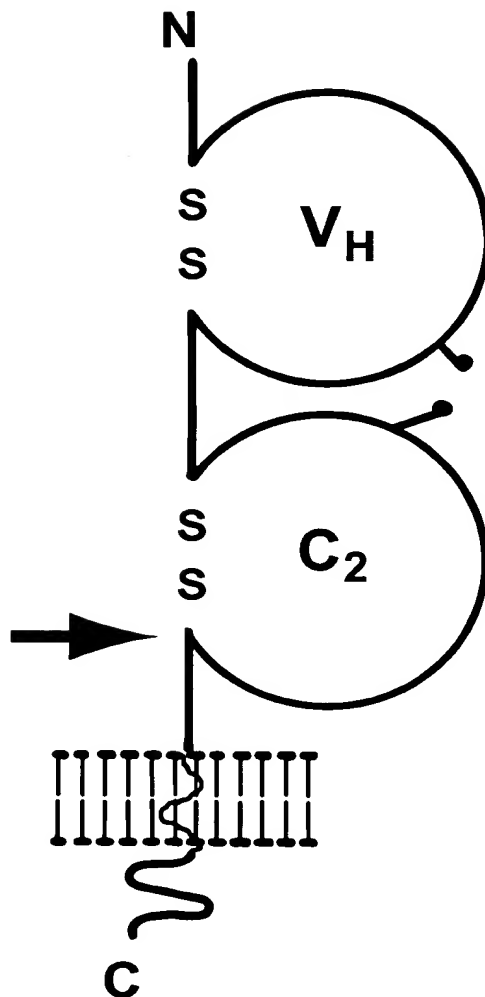


FIG. 8D



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FIG. 9A

APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

CRAM-1 :
CRAM-2 :
JAM :
Consensus :
20
MAISRRRLRLLYARLPHEFLFLFR
MARSPOGLMLLLHLLTVALDY
MGTEGKAGRKLLFLFTSMILG
r q l Llh f l
40

CRAM-1 :
CRAM-2 :
JAM :
Consensus :
60
GCMIEAVNLKSSNRNPVHFEFESVE
HKANGFSASKDHRQEVTVIEHQEAI
SLVQKGGSVYLAQSDVQVPEMESTK
q k V V Ef S

CRAM-1 :
CRAM-2 :
JAM :
Consensus :
80
* *
LS C I I T H S Q T S D P R I E W K K I O D G Q T
L A C K - T P K K T T S S R L E W K K V G Q G - V
L T C - - T Y S G F S S P R V E W K F V G S T T
L C T s t s s p R E W K K q g t
100

CRAM-1 : 100
CRAM-2 : 95
JAM : 93
Consensus :
LVY VYF D N K I O G D L A G R T D V F E G K T S L
S L V Y Y Q Q A L O G D F K D R A E M I D - F N I
A L V C Y N S Q I T A P Y A D R V - T F S S E G I i
lvv i o g d a d r f



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APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

FIG. 9B

* 120

RIWNVTRSDSAIYRCEVVALNDR-K
RIKNVTRSDAGEYRCEVSAPTQGO
TFSSVTRKDNGEYTCMVS--EEGGO
ri nVTRsd geYrCeVsa aa

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

140

*

EMDEITIELI VOVKPVTVCRI PAA
NEQEDKVMLEVLVAPAVPACVPTS
NYGEVSIHLTVLPVPSKPTISVPS
E L VLV P P C P S

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

160

*

VPVGKTATLQCOSEGYPRPHYSWY
VMTGSVVELRCODKFGNPAPEYIWF
VTIGNRAVLTCSEHDGSPPEYSWF
V G a L Cq G P peYSW

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

180

200

RNDVPLPT-DSRANPRFQNSSFHVN
KDGTSLLG-NPKGGTH-NNSSYTNE
KDGTSMLTADAKKTRAFHNSSFTID
k c s le k f NSS t

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

: 198
: 193
: 191



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FIG. 9C

APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

220*

SETGTLVFNAVHKDDSGQYYC IASN
HESGILQFNMISSKMDSGEYYC EARN
PKSGDLYFDFVTAFAFDSGEYYC QAOQN
esG L F k DSG YYC A N

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

240

*

DAGAA-RCEGQDM E V Y D L N I A G I I G
SVGHR-RCFPGKRM O V D V L N I S G I I A
GYGTAMRSEAAHMDAVE L N V G G I V A
G a Rceq M v L N G I a

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

260

GVLVLI V L A V I T M G I C C A Y R R G C F
TVVVVAFVISVCG L G T C M A O R K G Y F
AVLVTLLI L L G L I F G M F A Y S R G Y F
VlVvliVl v G c AvrrGvF

CRAM-1 :
CRAM-2 :
JAM :
Consensus :

280

300

ISSKQDGESYKSPGKHDGVNYLRTS : 297
--SKETS FQKGS PASK - - - - VTMT : 285
--ETTKKGTA PGKKVIYSQPSTR : 287
sk t sPqkk T

CRAM-1 :
CRAM-2 :
JAM :
Consensus :



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APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 9D

CRAM-1	:	EEGD	FRHK	SS	FF	VI	:	310
CRAM-2	:	GEN	FRHT	KS	FF	II	:	298
JAM	:	SEGE	FKOT	SS	FF	LV	:	300
Consensus	:	Eq	Frht	ss	FF			



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APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

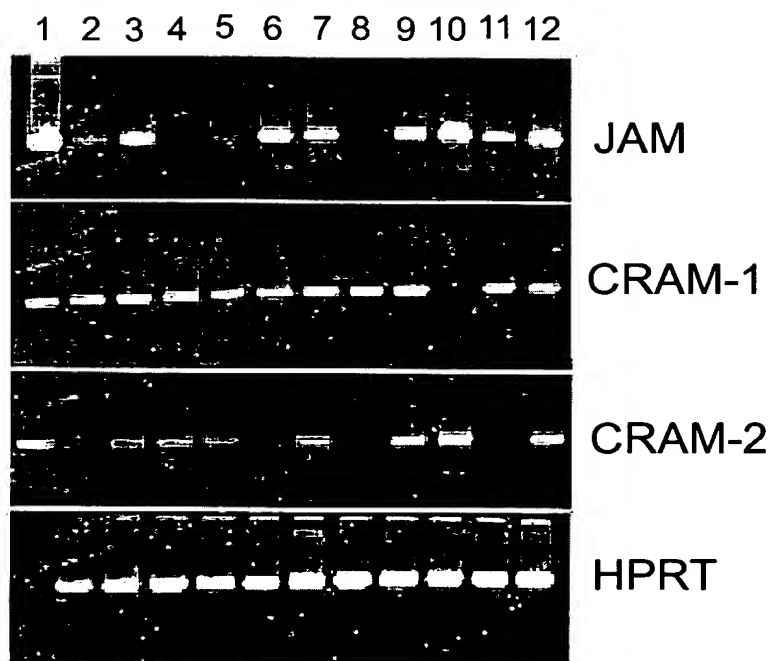


FIG. 10A



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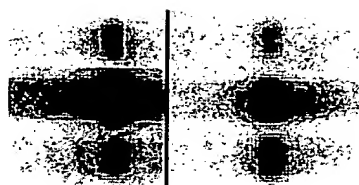
FIG. 10B

APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
DRAFTSMAN		

Lymph Nodes
Peyer's Patches
Thymus

JAM

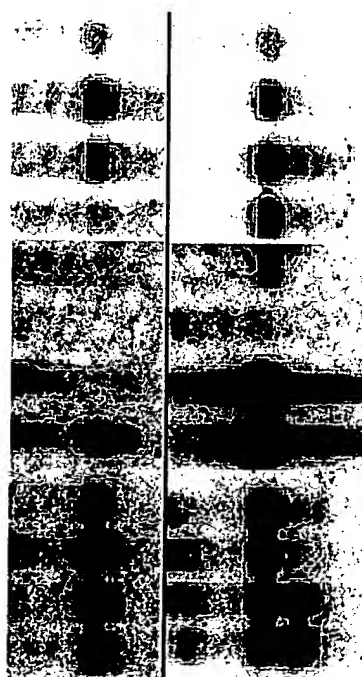
β -ACTIN



Intestine
Lymph Nodes
Peyer's Patches
Thymus
Liver
Placenta
KLN205
TME
E17
E15
E11
E7

CRAM-1

β -ACTIN





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O.G. FIG.		SUBCLASS
APPROVED	BY	CLASS
DRAFTSMAN		

FIG. 11A

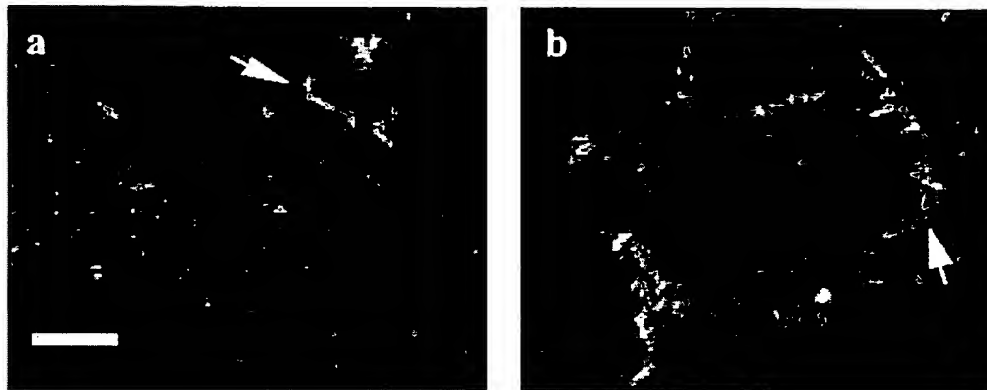
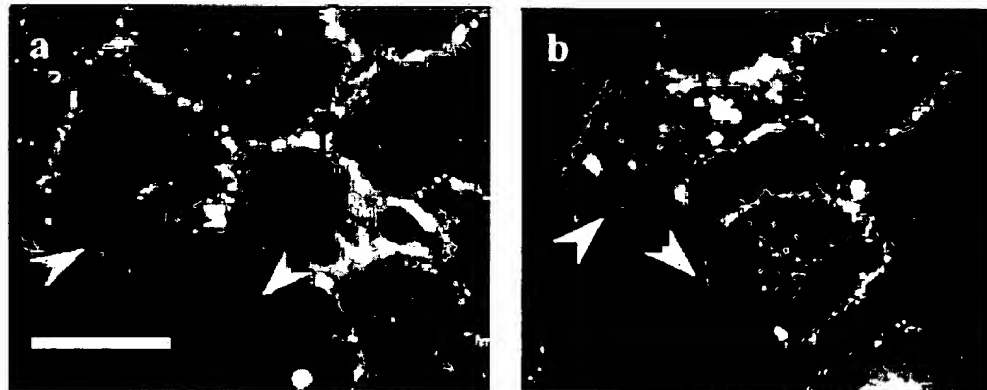


FIG. 11B





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APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

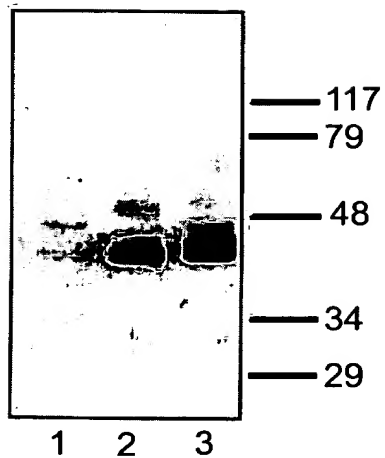


FIG. 11C

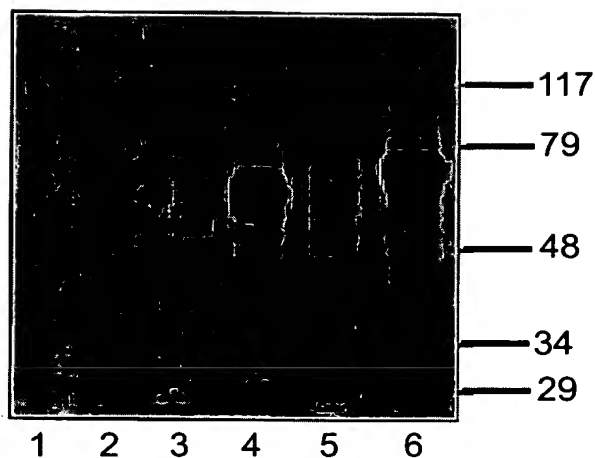


FIG. 11D



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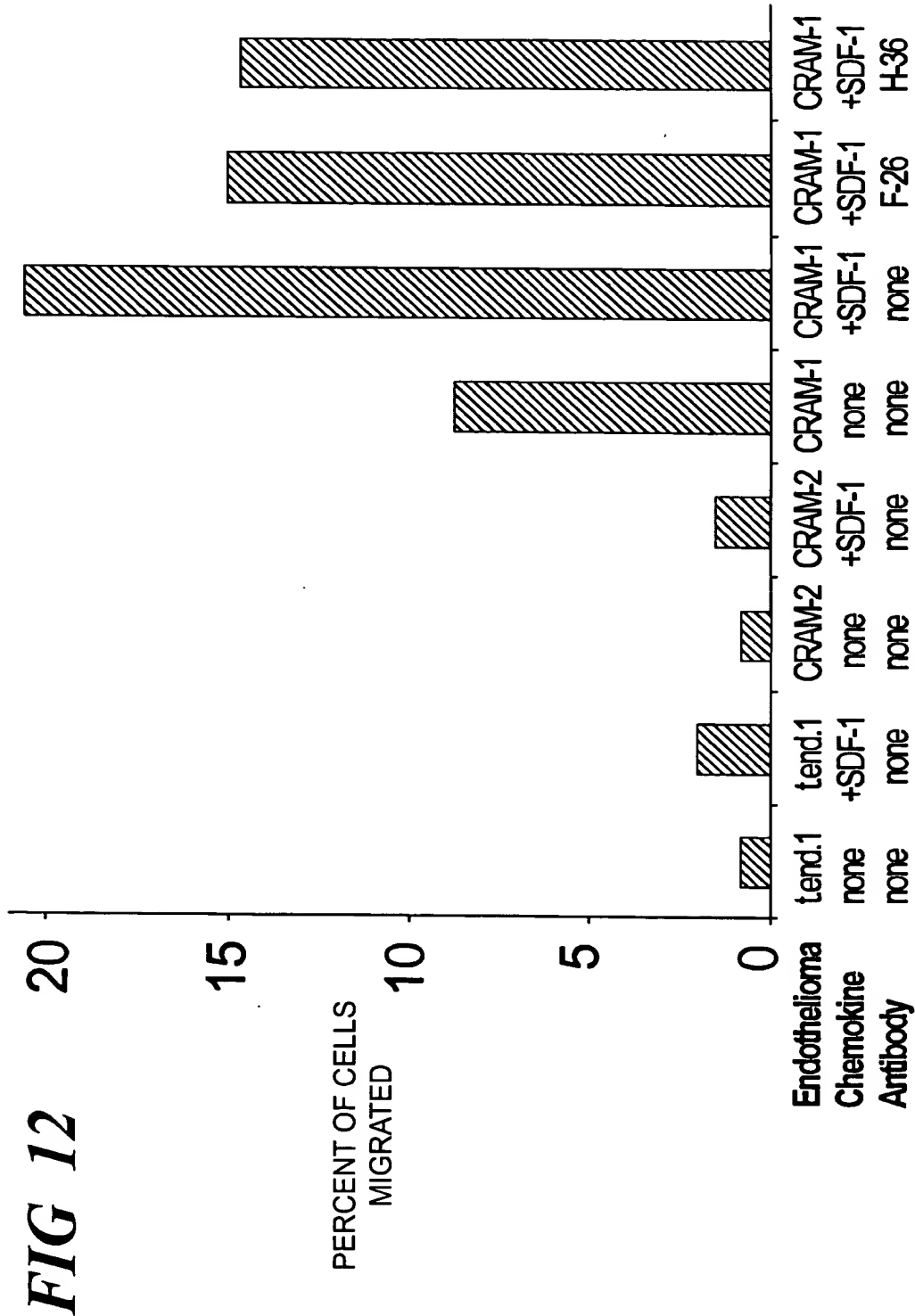
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APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		



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O.G. FIG.	
CLASS	SUBCLASS
APPROVED BY DRAFTSMAN	

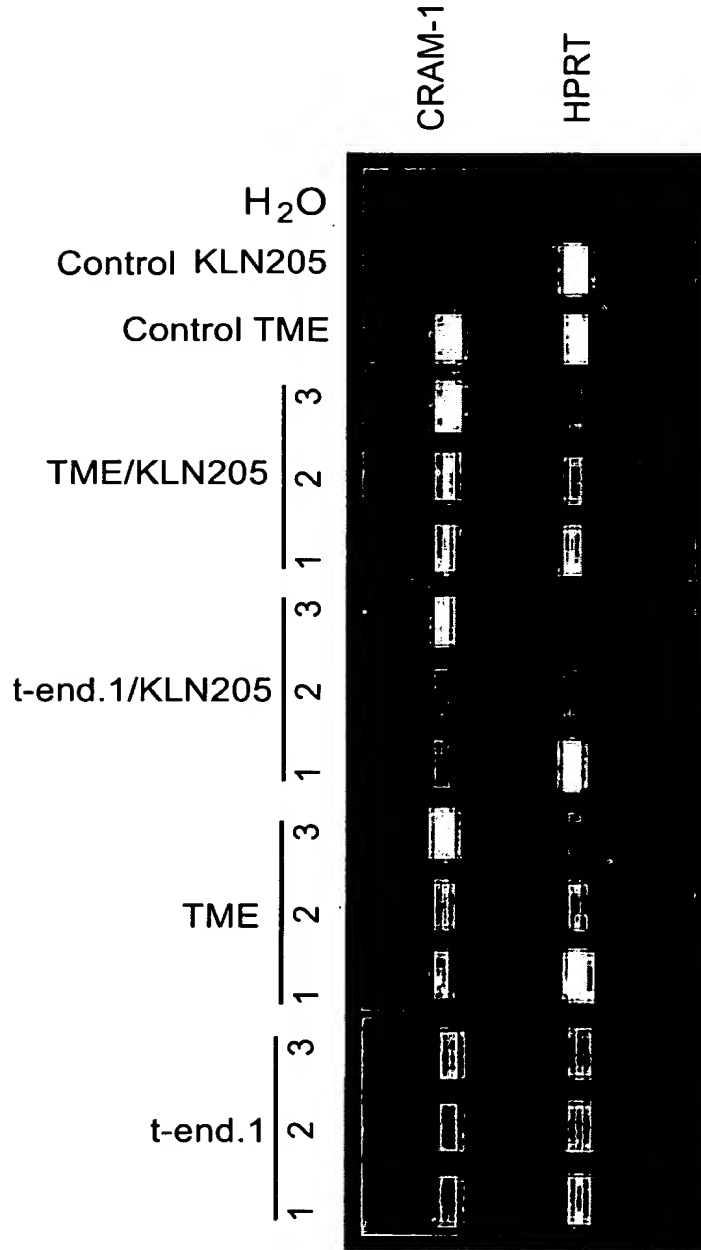


FIG. 13



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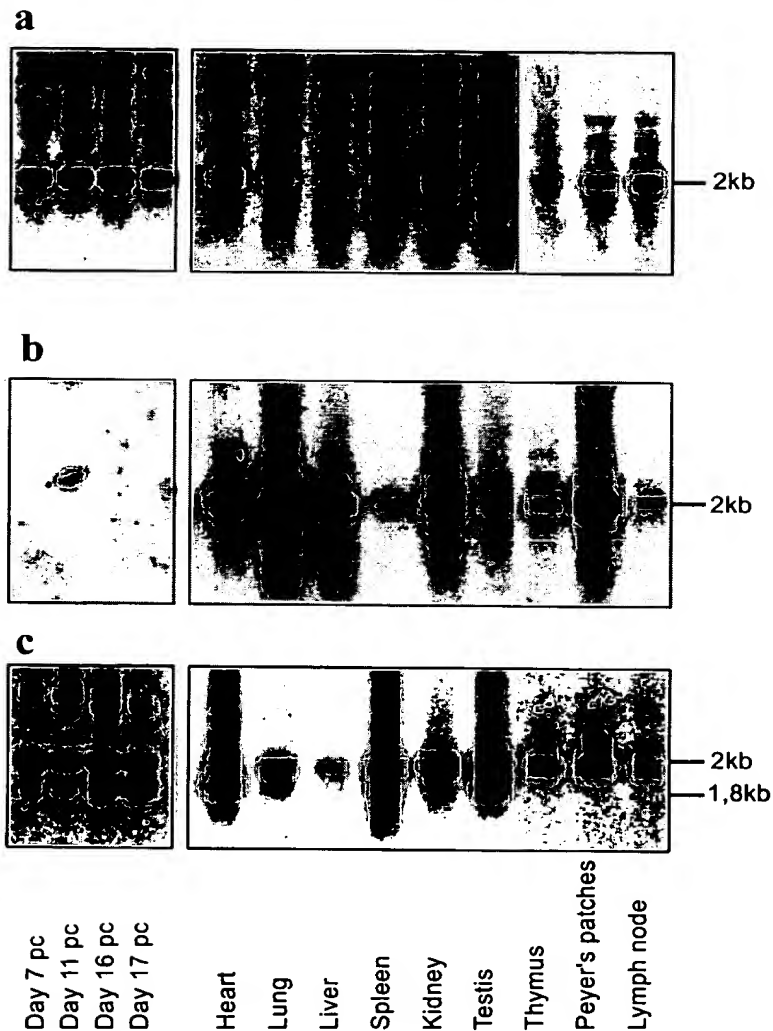
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APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

FIG. 14





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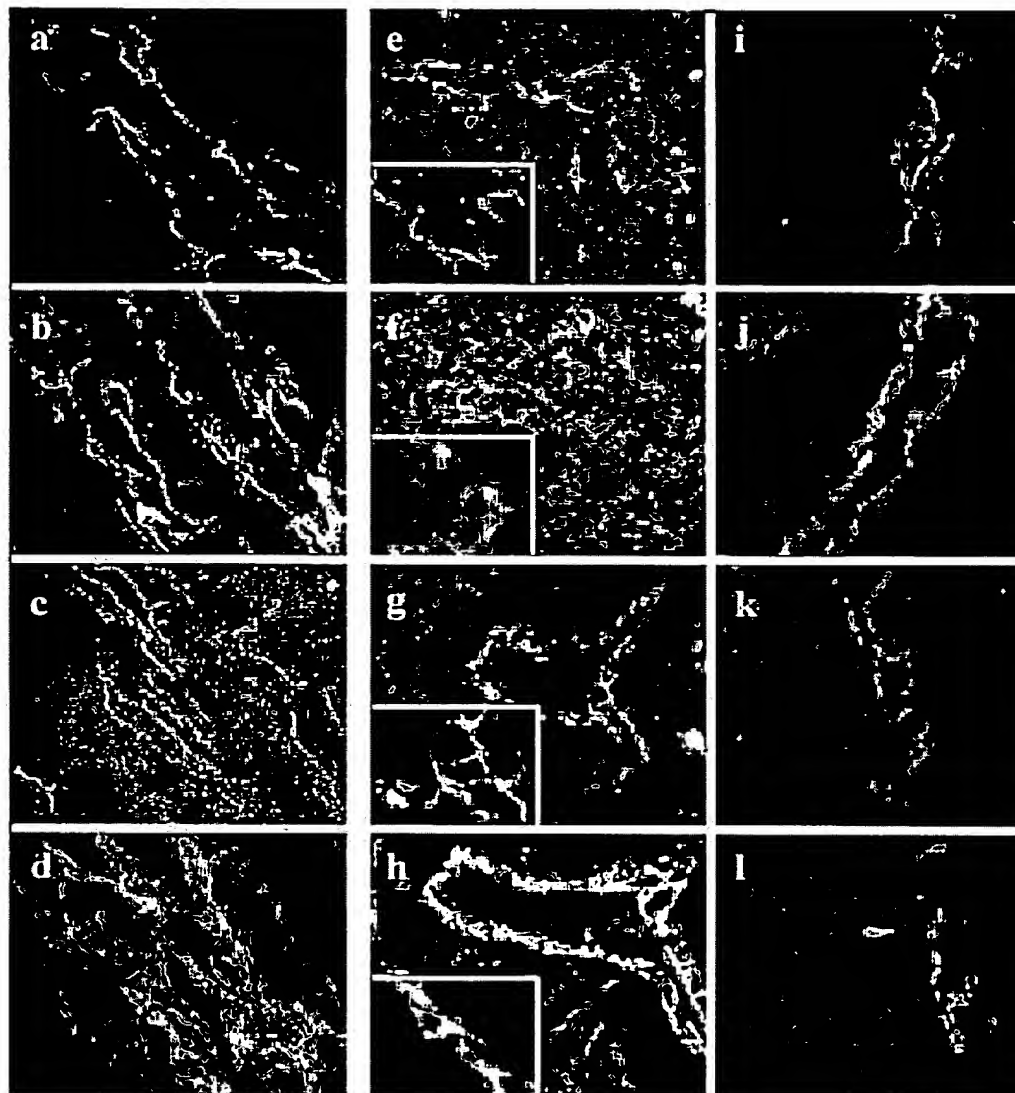
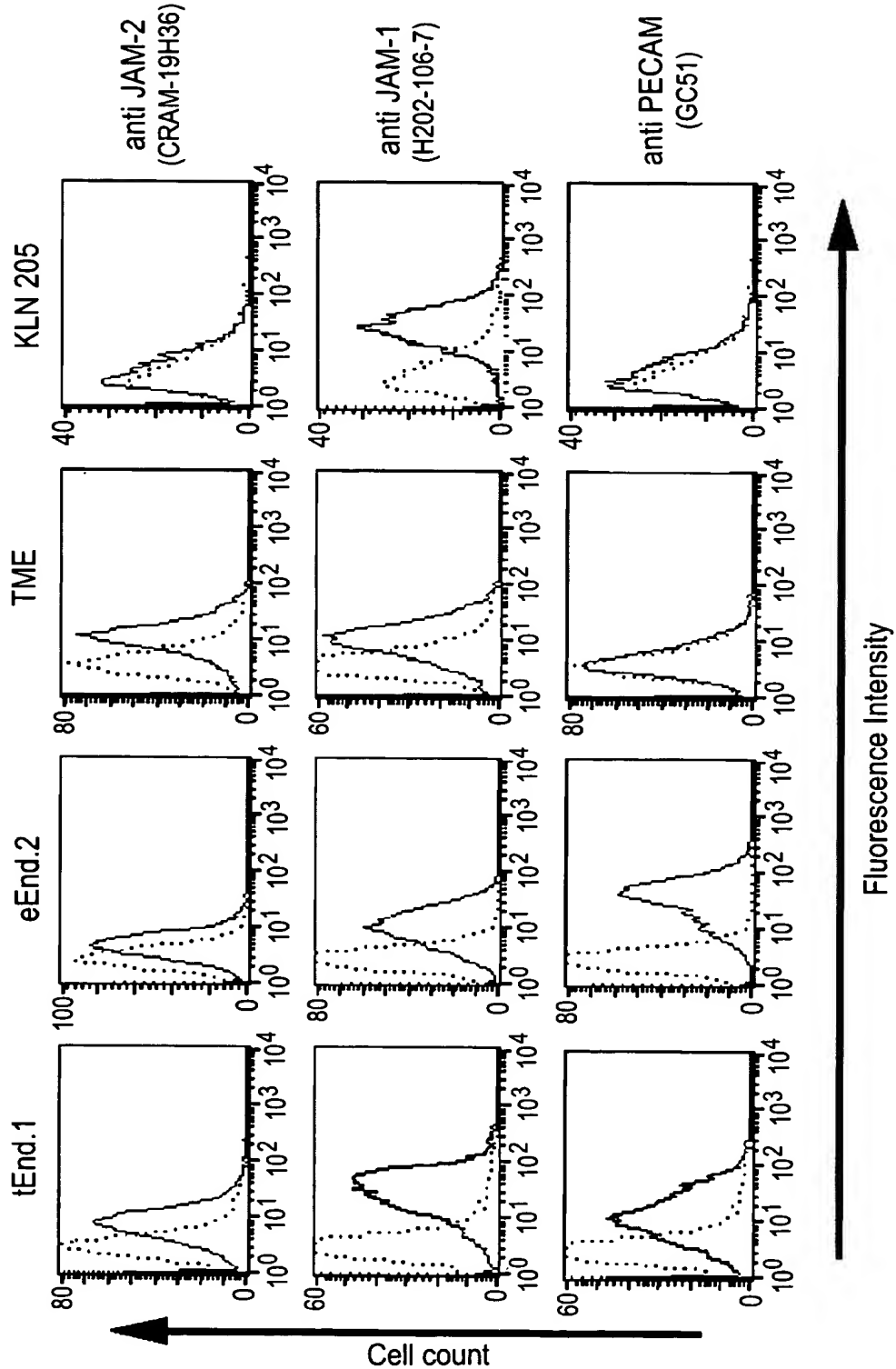


FIG. 15

O.G. FIG.	
APPROVED	BY
CLASS	SUBCLASS
DRAFTSMAN	

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 16A



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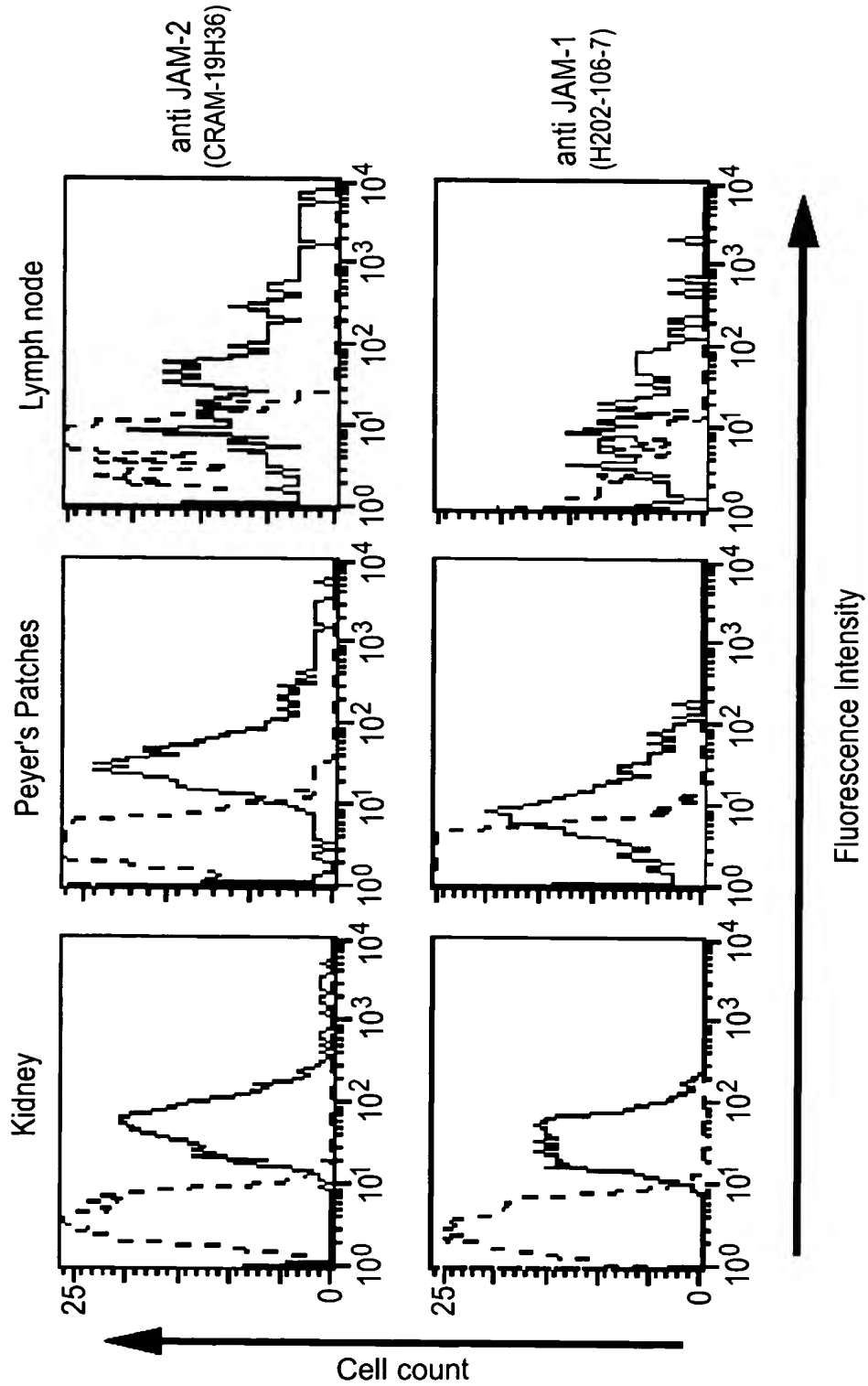
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FIG. 16B



APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
CRAFTSMAN		





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APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

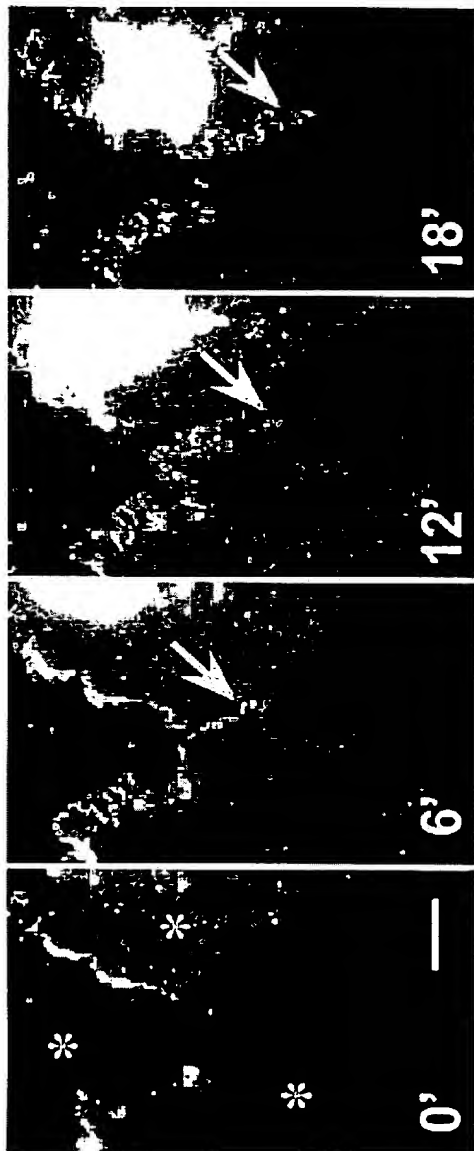


FIG. 17A

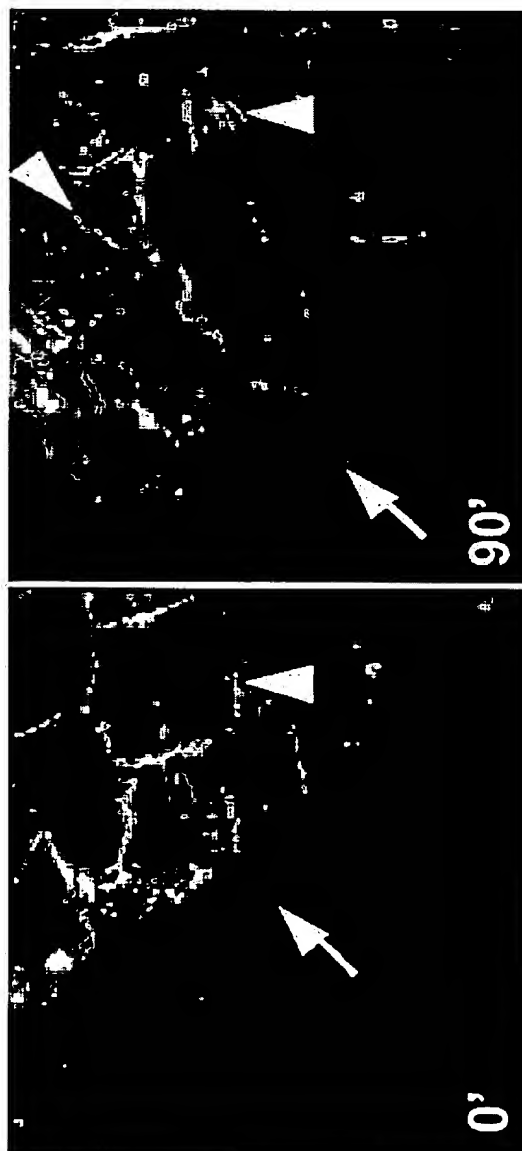


FIG. 17B



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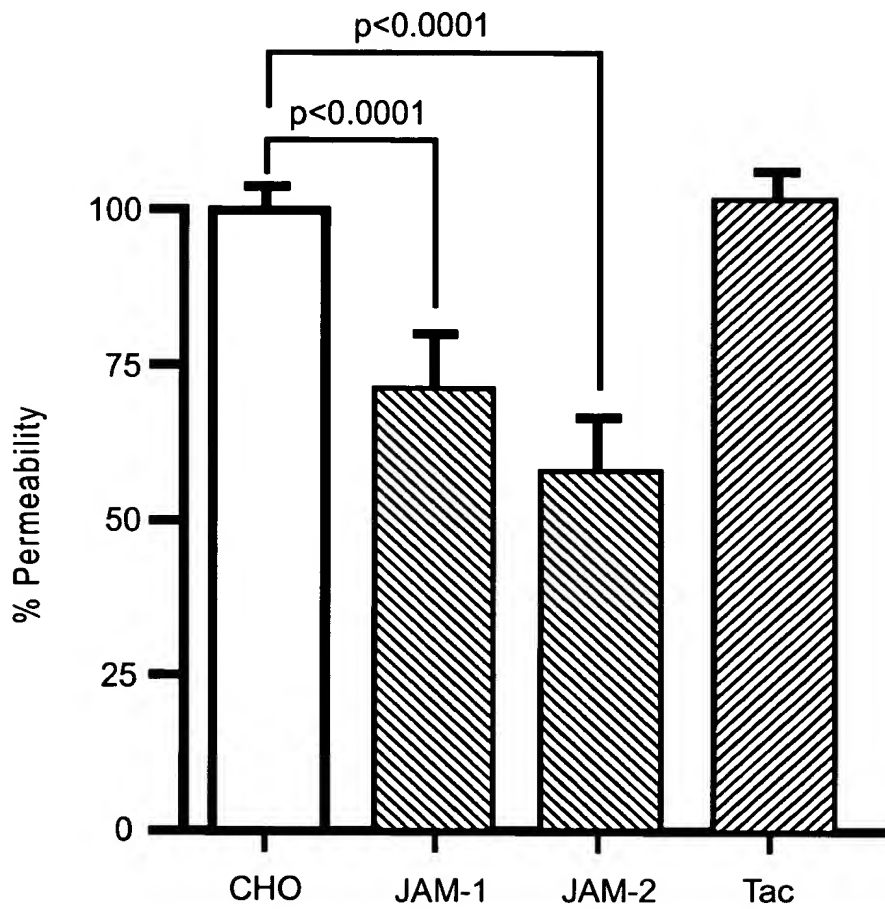


FIG. 18

APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
DRAFTSMAN		



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APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
DRAFTSMAN		

Basal → Apical

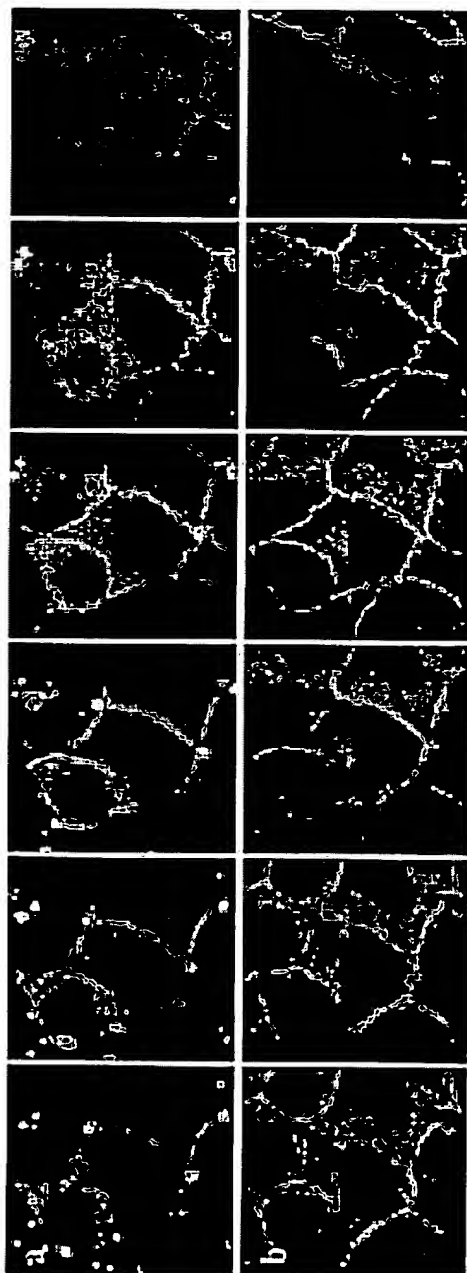


FIG. 19A

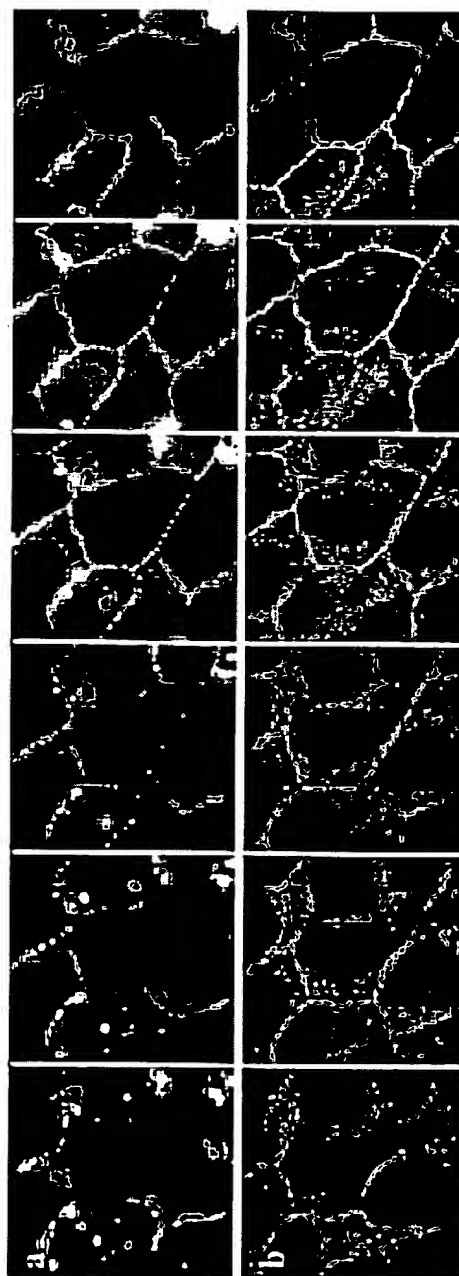


FIG. 19B



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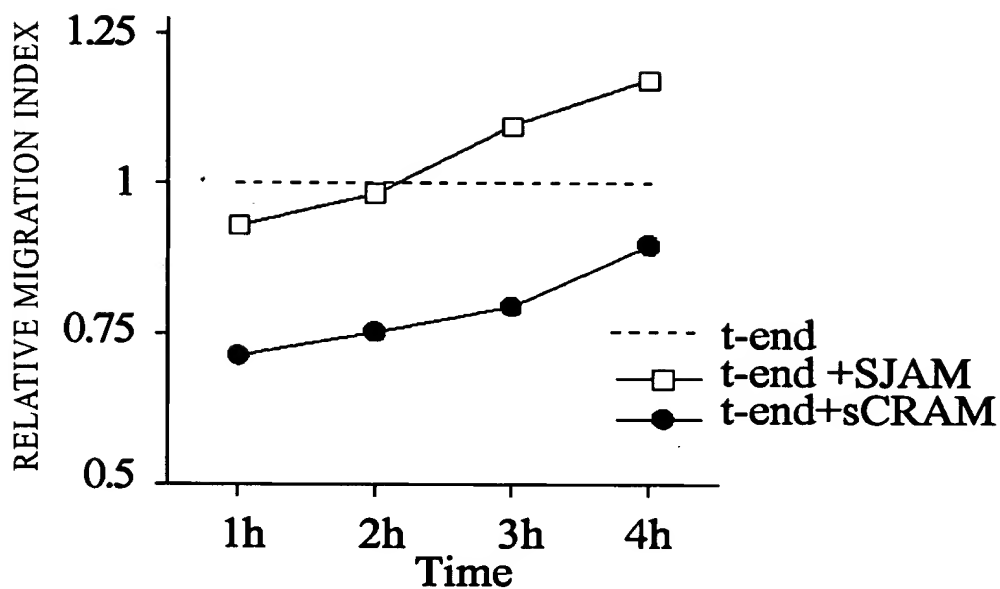
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FIG. 20A



APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
DRAFTSMAN		



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APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 20B

